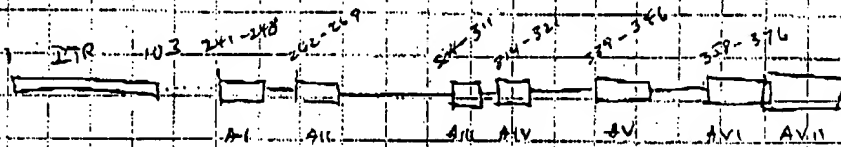


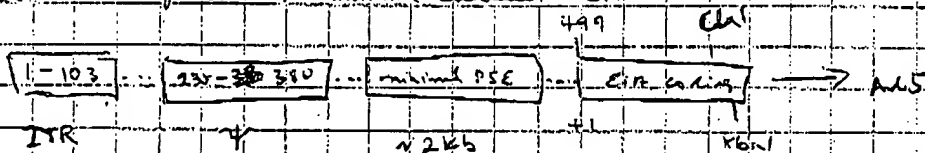
**BEST AVAILABLE COPY**

In designing the protein-specific AAS (AAS PSE) it did already work on CIA enhancer and for  $\psi$  signal (packaging). There are 6 A-rich sequences for  $\psi$  and they can be moved to the left up against the ITR on left end of the genome.



possible element I of evidence at 199-205 can be deleted.

Link PCR + ITR - V - minimal enhancer - EIA



$$\text{Deposits of A/S } 235-103 = 132$$

$$490 - 380 = 110$$

but add back ~ 2 kb, making 20 kb for minimal PSE

$\therefore$  total add 1.8 kb

Do this by PCR replacing wt in exc

~~SECRET~~ 2

p BTG-11 days  $\sim 3kb$  in  $\Delta E3$

$3.0 - 1.8 = 1.2$  to use for residual safety or confinement.

also 5% compression  $57 + (0.05) \times 35 \text{ kPa} = 1750 \text{ kPa}$

$$\therefore \text{absolute max} = 1.2 + 1.7 \text{ kg} = 2.9$$

∴ use a smaller enhancer for HSV-Ek on cytosine deaminase.

→ This should be minimal PSE of elements responsive to EIA.

To Page No. \_\_\_\_\_

Witnessed & Understood by me,

Date \_\_\_\_\_

**Invented by**

Date \_\_\_\_\_

**R corded by**

# BEST AVAILABLE COPY

Did minis 12 on 8 plate 8, 12, 14, 15-23.  
What should they be cut with?

1-12 from plate 8: KpnI & ClaI

13-24 from plate 12: ClaI

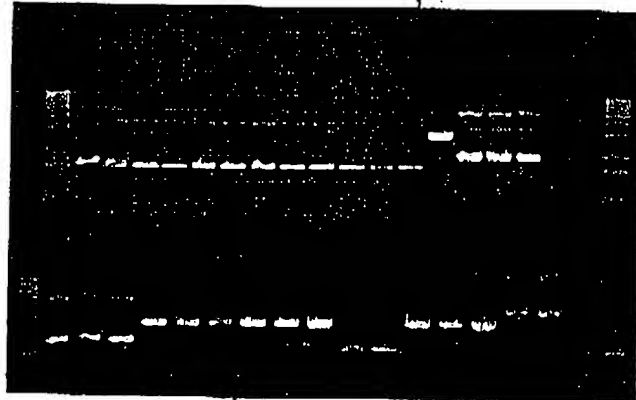
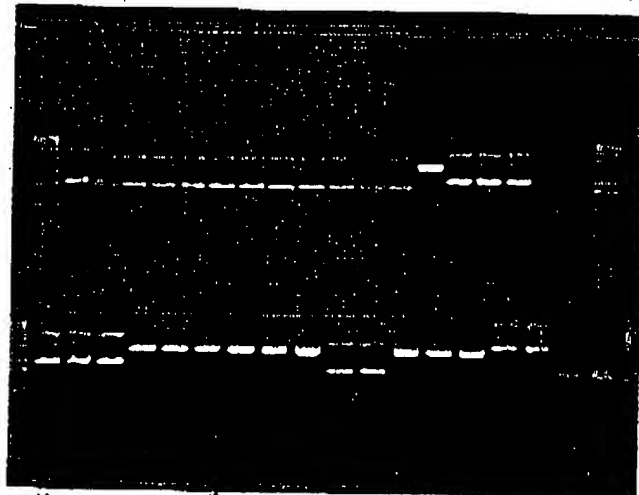
25-26 from plate 16: ClaI

all w/ ClaI

1-12

1 12-18

use 4 µl DNA



2 µl from  
Microbiol. com. 1  
So not 203 pages 25  
except no dig in. Empty

To Page No. \_\_\_\_\_

Witnessed & Understood by me,

*J. Anderson*

Invented by

Date

Recorded by

*J. Anderson*